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The Genetic Architecture of Arsenic Metabolism Efficiency: A SNP-Based Heritability Study of Bangladeshi Adults

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LMM Analysis

Reference

Figure S1. Regional heritability estimates (A) and corresponding P-values (B) for DMA% based on all samples (n= 2,053). Estimates were obtained using imputed SNP data with window size 300 SNPs with a 50 SNP overlap between windows. 4,787 tests were conducted. Red line indicates Bonferroni-corrected p threshold.